ENTERED



RAW SEQUENCE LISTING

1 <110> APPLICANT: Hangauer Jr., David G.

Marsilje, Thomas H.

PATENT APPLICATION: US/09/482,585A

DATE: 11/22/2002 TIME: 11:03:51

Input Set : N:\Crf4\11182002\I482585.raw
Output Set: N:\CRF4\11222002\I482585A.raw

```
Milkiewicz, Karen L.
      3
      4 <120> TITLE OF INVENTION: A NOVEL METHOD FOR DESIGNING PROTEIN KINASE INHIBITORS
      5 <130> FILE REFERENCE: 19226/931
      6 <140> CURRENT APPLICATION NUMBER: US/09/482,585A
      7 <141> CURRENT FILING DATE: 2000-01-13
      8 <150> PRIOR APPLICATION NUMBER: 60/115,643
      9 <151> PRIOR FILING DATE: 1999-01-13
     10 <160> NUMBER OF SEQ ID NOS: 7
     11 <170> SOFTWARE: PatentIn Ver. 2.1
     13 <210> SEQ ID NO: 1
     14 <211> LENGTH: 5
     15 <212> TYPE: PRT
     16 <213> ORGANISM: Artificial Sequence
     17 <220> FEATURE:
     18 <223> OTHER INFORMATION: Description of Artificial Sequence: src substrate
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     22
               1
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     26 <212> TYPE: PRT
     27 <213> ORGANISM: Artificial Sequence
     28 <220> FEATURE:
     29 <221> NAME/KEY: PEPTIDE
     30 <222> LOCATION: (2)
     31 <223> OTHER INFORMATION: Xaa in position 2 is modified Tyr.
     32 <220> FEATURE:
     33 <223> OTHER INFORMATION: Description of Artificial Sequence: src
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W-->36
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               1
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     40 <211> LENGTH: 5
     41 <212> TYPE: PRT
     42 <213> ORGANISM: Artificial Sequence
     43 <220> FEATURE:
     44 <221> NAME/KEY: PEPTIDE
     45 <222> LOCATION: (4)
     46 <223> OTHER INFORMATION: Xaa in position 4 is modified Ala.
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DATE: 11/22/2002

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PATENT APPLICATION: US/09/482,585A
                                                               TIME: 11:03:51
                     Input Set : N:\Crf4\11182002\I482585.raw
                     Output Set: N:\CRF4\11222002\I482585A.raw
     47 <220> FEATURE:
     48 <223> OTHER INFORMATION: Description of Artificial Sequence: PKA
              pentapeptide scaffold
     50 <400> SEQUENCE: 3
W--> 51
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     52
     54 <210> SEQ ID NO: 4
     55 <211> LENGTH: 5
     56 <212> TYPE: PRT
     57 <213> ORGANISM: Artificial Sequence
     58 <220> FEATURE:
     59 <221> NAME/KEY: PEPTIDE
     60 <222> LOCATION: (4)
     61 <223> OTHER INFORMATION: Xaa in position 4 is Ala or modified Ala.
     62 <220> FEATURE:
     63 <223> OTHER INFORMATION: Description of Artificial Sequence: Boronic
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     65 <400> SEQUENCE: 4
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     69 <210> SEQ ID NO: 5
     70 <211> LENGTH: 7
     71 <212> TYPE: PRT
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     73 <220> FEATURE:
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     76
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     77
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     85 <222> LOCATION: (5)
     86 <223> OTHER INFORMATION: Xaa in position 5 is ALA; PHOSPHORYLATION
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     94 <210> SEQ ID NO: 7
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     96 <212> TYPE: PRT
     97 <213> ORGANISM: Artificial Sequence
     98 <220> FEATURE:
     99 <223> OTHER INFORMATION: Description of Artificial Sequence: Peptide
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RAW SEQUENCE LISTING

RAW SEQUENCE LISTING

DATE: 11/22/2002 PATENT APPLICATION: US/09/482,585A TIME: 11:03:52

Input Set : N:\Crf4\11182002\I482585.raw Output Set: N:\CRF4\11222002\I482585A.raw

substrate for Src 100

101 <400> SEQUENCE: 7

Gly Ile Tyr Trp His His Tyr

103

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 11/22/2002 PATENT APPLICATION: US/09/482,585A TIME: 11:03:53

Input Set : N:\Crf4\11182002\I482585.raw
Output Set: N:\CRF4\11222002\I482585A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 2
Seq#:3; Xaa Pos. 4
Seq#:4; Xaa Pos. 4
Seq#:6; Xaa Pos. 5

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 4

VERIFICATION SUMMARY

DATE: 11/22/2002 TIME: 11:03:53 PATENT APPLICATION: US/09/482,585A

Input Set : N:\Crf4\11182002\I482585.raw Output Set: N:\CRF4\11222002\I482585A.raw

L:36 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0 L:66 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0 L:91 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0